

Changes in Immune Cell Distribution and IL-10 Production are Regulated through Endometrial IP-10 Expression in the Goat Uterus

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PROBLEM: Changes in distribution or redistribution of immune cells are required for the establishment and maintenance of pregnancy, but these changes during early pregnancy have been poorly understood in the ruminant ungulates. Expression of a chemokine, interferon- γ (IFN- γ)-inducible protein 10 kDa (IP-10, CXCL10), was identified in the endometrium of pregnant goats. Population and/or distribution of endometrial immune cells and their cytokine productions could be regulated by IP-10 during the period of pregnancy establishment.

METHOD OF STUDY: Using reverse transcriptase-polymerase chain reaction (RT-PCR), expression of IP-10, IFN- γ , tumor necrosis factor- α , interleukin-10 (IL-10), CXCR3 mRNA and leukocyte cell surface markers, CD4, CD8, CD11b and CD45 mRNA during the caprine early pregnancy was investigated. The ability of IP-10 to stimulate peripheral blood mononuclear cells (PBMCs) migration was demonstrated using a chemotaxis assay. Changes in migration of PBMCs' immune cell population and cytokine expressions with IP-10 stimulation were investigated using flow cytometry and RT-PCR respectively.

RESULTS: Levels of IP-10, IL-10, CD4 and CD11b mRNA, and the number of CD4 and CD11b positive cells in pregnant goat endometrium were higher than those of cyclic goat endometrium. Migration of PBMCs was stimulated by recombinant caprine IP-10, and the effect was significantly reduced by neutralization with the use of an anti-IP-10 antibody. In the flow cytometric and RT-PCR analyses, migrated cells stimulated by IP-10 increased the expression of IL-10 and CD11b mRNA. Furthermore, IP-10 could stimulate the expression of IL-10 mRNA from PBMCs.

CONCLUSION: Endometrial chemokine IP-10 could regulate IL-10 production by resident and possibly migrated cells expressing CD11b, probably natural killer cells, and these changes may result in immune environments of the uterus suitable for conceptus implantation in ruminants.

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INTRODUCTION

In addition to biochemical communications, an interaction between the conceptus and maternal immune system may also be important in the establishment of

pregnancy. Maternal immune system is highly complex and must be tightly regulated to elicit an effective and appropriate response to the conceptus. In ruminant ungulates, high incidences of conceptus losses, which may result from insufficient cellular and biochemical

interactions, occur during early pregnancy.^{1,2} Such losses may also result from a lack of suitable changes in local immune environments, i.e. types and number of immune cells at or near implantation sites. Although exact mechanisms by which the maternal immune environments are established during the period of early pregnancy are not fully understood, it is likely that changes in the distribution of immune cells³ and patterns of local cytokine productions are involved in the establishment of proper microenvironments *in utero*.⁴⁻⁷

Cytokines are soluble factors functioning in the immune systems, and acting as mediators for communication networks between cells of both lymphoid and non-lymphoid origins. Such molecules exert pleiotropic effects on these cells such as regulation of migration, activation, proliferation and possibly antibody production. These cytokines are functionally divided into proinflammatory cytokines, such as tumor necrosis factor- α (TNF- α) and interferon- γ (IFN- γ), and regulatory cytokines such as interleukin-10 (IL-10) and transforming growth factor- β . It is believed that the local immunoregulation during early pregnancy is biased toward the increased production of regulatory cytokines, which suppress the proinflammatory reactions.^{8,9} One of regulatory cytokines, IL-10, is thought to be produced by T helper 2 (Th2) lymphocytes; however, recent studies have shown that natural killer (NK) cells are also capable of producing IL-10.^{10,11} In humans and mice, the redistribution of immune cells during early pregnancy is considered prerequisite for successful implantation, particularly increases in the number of uterine NK (uNK) cells at the implantation site.^{12,13} These uNK cells are thought to produce IL-10, which in turn alters uterine environments suitable for embryonic survival. In ruminant ungulates, NK-like cells with lytic activity were found in the ovine endometrium during early pregnancy.¹⁴ It is therefore suspected that uNK or NK-like cells also play an important role during the period of implantation in ruminants. However, biochemical and molecular mechanisms by which these cells migrate and/or are recruited to an implantation site have not been well characterized.

In ruminants, a major protein, IFN- τ , produced by the developing blastocyst is implicated in the process of maternal recognition of pregnancy.¹⁵⁻¹⁷ IFN- τ prevents the regression of corpus luteum (CL), at least in part by inhibiting endometrial estrogen receptor expression, thus preventing estrogen stimulation of the oxytocin receptor, which impedes pulsatile releases of endometrial luteolysin, prostaglandin F_{2 α} .¹⁸ Furthermore, IFN- τ is known to regulate lymphocyte proliferation¹⁹⁻²¹ and cytokine production,^{22,23} suggesting that IFN- τ may also play a role in the changes and/or adjustment of immunological environments

suitable for the process of implantation. Recently, the endometrial expression of a chemokine, IFN- γ -inducible protein 10 kDa (IP-10, CXCL10), during the period of sheep and goat implantation was found in our laboratory. More importantly, maternal expression of IP-10 was stimulated by conceptus IFN- τ .^{24,25} Chemokines are a family of structurally related cytokines with low molecular masses and play a key role in multiple aspects of inflammatory and immune responses by promoting migration and activation of different subpopulations of immune cells.²⁶ IP-10 is a member of the C-X-C chemokines and targets NK cells and Th1 lymphocytes through the C-X-C chemokine receptor 3 (CXCR3).^{27,28} Because other chemokines are also expressed in the endometrium of ruminant ungulates,^{29,30} it is suspected that local expressions of chemokines, which play a role in the distribution or redistribution of immune cells and their cytokine productions, are required for successful pregnancy.

The present study was undertaken to gain better insight into the distribution and/or redistribution of immune cells and subsequent changes in cytokine productions by one of endometrial chemokines, IP-10, in the goat uterus. Transcript changes in IP-10, IFN- γ , TNF- α , IL-10, CXCR3 and leukocytes' cell surface markers, CD4, CD8, CD11b and CD45 were studied during early pregnancy. Studies were extended to examine whether IP-10 could regulate the migration of peripheral blood mononuclear cells (PBMCs) and to investigate changes in cell phenotypes and/or expression of cytokine genes in the migrated PBMCs.

MATERIAL AND METHODS

Animals and Tissue Preparation

Shiba goats were maintained at the University of Tokyo farm and the protocol for goat experimentation has been reviewed and approved by the animal care committee at the University. Uteri from cyclic goats on day 15 ($n = 7$) and pregnant goats on days 17 ($n = 3$) and 20 ($n = 5$) were removed by hysterectomy under isoflurane anesthesia.²⁵ Two uteri each from day 15 cyclic goats and day 20 pregnant goats were frozen immediately for subsequent immunohistochemical studies. Endometrial tissues collected from goats ($n = 3$ for each day examined) were frozen immediately and stored at -70°C ; these frozen tissues were subjected to RNA extraction. Whole blood was collected from day 15 cyclic goats, and PBMCs were used for IP-10 stimulation and chemotaxis assays.²⁵

In Vitro Culture

The PBMCs were separated from ethylenediaminetetraacetic acid-treated whole blood (80 mL) by

density gradient centrifugation ($800 \times g$ at 20°C for 30 min, OptiPrep; Nycomed, Roskilde, Denmark) and were suspended in RPMI 1640 medium supplemented with 10% fetal calf serum (Sigma, St Louis, MO, USA), 40 U/mL of penicillin, 40 $\mu\text{g/mL}$ of streptomycin and anti-pleuropneumonia-like organisms agent (Invitrogen Corp., Carlsbad, CA, USA). PBMCs counted and adjusted to 3×10^7 cells/mL were plated onto 6-well costar plates (3 mL per well), which were then incubated with 20 ng/mL recombinant caprine IP-10 (rcIP-10),²⁵ 10 ng/mL recombinant bovine IFN- τ (Katakura Industries Co., Tokyo, Japan) or control (20 ng/mL recombinant glutathione S-transferase, GST)²⁵ at 37°C , 5% CO_2 and 95% air atmosphere for 8 hr. After incubation, cells were immediately processed for total RNA extraction.

Chemotaxis Assay

Migration of PBMCs was assessed in a 96-well modified Boyden chamber (NeuroProbe, Cabin John, MD, USA) using polyvinylpyrrolidone-free polycarbonate membrane (5 μm pore size, NeuroProbe), as described previously.²⁵ In brief, isolated PBMCs were labeled with the intracellular fluorescent dye, calcein-AM (4 μM , Molecular Probes Inc., Eugene, OR, USA), at 37°C for 30 min. Dulbecco's modified Eagle's medium (DMEM)-0.1% bovine serum albumin (BSA) (without phenol red) supplemented with 20 ng/mL rcIP-10 was added to the bottom wells of the chemotaxis chamber, whereas labeled PBMCs (5×10^6 cells/mL) in DMEM-0.1% BSA (without phenol red) were added to the top wells of the chamber. After the chambers were incubated at 37°C and 5% CO_2 atmosphere for 2 hr, fluorescence of cells in the bottom wells was measured using a fluorescence reader (excitation filter 485 nm and emission filter 535 nm, ARVO SX 1420 Multilabel Counter, Perkin-Elmer Life Sciences Inc., Boston, MA, USA). For the blocking experiments, rcIP-10 was pre-incubated at 37°C for 1 hr with 30 $\mu\text{g/mL}$ of a rabbit polyclonal antibody to rcIP-10 or control rabbit IgG (Sigma) before addition to the bottom chamber.²⁵ Three replicates were performed for each treatment, and three independent experiments were carried out for each treatment.

Flow Cytometric Analysis

To determine cell phenotypes, migrated cells on the chemotaxis assay were subjected to flow cytometric analysis (fluorescence-activated cell sorter, FACS). Chemotaxis assay of non-fluorescent-labeled PBMCs was performed as aforementioned.²⁵ Migrated cells were collected and blocked with Block Ace (Dainippon Pharmaceutical, Osaka, Japan) at 4°C for 30 min to prevent non-specific binding of antibodies, and then

incubated with 1:10 dilution of monoclonal antibodies to CD4, CD8 (generously provided by Dr Barcham, The University of Melbourne) or CD11b (Serotec Inc., Raleigh, NC, USA) at 4°C for 1 hr. After the incubation, cells were washed in cold phosphate-buffered saline, and incubated with fluorescein isothiocyanate (FITC)-conjugated goat anti-mouse IgG (15 $\mu\text{g/mL}$, Jackson ImmunoResearch Laboratories Inc., West Grove, PA, USA) at 4°C for 30 min. Flow cytometric analyses were performed using a FACS Calibur cytofluorimeter (Becton Dickinson, Mountain View, CA, USA).

Immunohistochemical Analysis

Frozen whole uteri sections (10 μm) were mounted onto silane-coated slides and fixed in acetone. Non-specific binding was blocked by 1-hr treatment with Block Ace at room temperature, followed by incubation with monoclonal antibodies to CD4, CD8 or CD11b at 4°C for 12 hr. Slides were then incubated with biotin-conjugated donkey anti-mouse IgG (1:400, Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA) at room temperature for 1 hr. Immunoreactions were visualized by diaminobenzidine (Sigma). An equal concentration of normal mouse IgG was used on serial sections of the same tissue to evaluate non-specific binding (negative control). Tissue sections were then counterstained with hematoxylin.

RT-PCR Analysis

Using Isogen (Nippon Gene, Tokyo, Japan), total RNAs were extracted from endometrial tissues and PBMCs.²⁴ Total RNA samples (2 μg) were first reverse transcribed with SuperScriptII (Invitrogen) and oligo-dT primers (20 μL reaction volume).²⁵ Relative mRNA levels of IP-10, CXCR3, IL-10, IFN- γ , TNF- α , CD4, CD8, CD11b, CD45 and glyceraldehyde 3-phosphate dehydrogenase (G3PDH) were determined from PCR amplification using oligonucleotide primers (Table I). Each reaction consisting of primer pairs was run with RT template (1 μL) and AmpliTaq Gold (0.625 U; Roche Molecular Systems, Branchburg, NJ, USA) in a final volume of 25 μL . All PCR reactions consisted of 35 or 40 cycles at 95°C for 1 min, 60°C for 1 min and 72°C for 1 min, followed by the final extension at 72°C for 12 min. Following agarose gel electrophoresis and visualization with ethidium bromide, PCR products were analyzed using an image analysis system (ATTO Corporation, Tokyo, Japan) equipped with the Quantity One (v3.0 software; PDI, Inc., Huntington Station, NY, USA). The identity of each amplified PCR product was verified by sequence analysis using a Perkin-Elmer sequencer (model ABI Prism 377 XL, Roche) after cloning into the pGEM-T Easy vector (Promega, Madison, WI, USA).

TABLE I. Oligonucleotide Primers Used for PCR Reactions

Name	Sequence of forward and reverse primer	No. of cycles	Product size (bp)
IP-10	5'-CAAGGCATACCTCTCTCTAG-3' 5'-TTCAGACATCTTTTCTCCCC-3'	40	170
CXCR3	5'-GCATCAGCTTCGATCGGTAC-3' 5'-GATGCGGGCGTAGCAATAGG-3'	40	284
IL-10	5'-CTGTCTGACAGCAGCTGTAC-3' 5'-ATACGTACAGAGAAGCTCA-3'	40	646
IFN- γ	5'-CGATGAAATACACAAGCTCC-3' 5'-GATTACATTGATGCTCTCCG-3'	40	504
TNF- α	5'-ACAACGGGCCACCAACCATC-3' 5'-TCTCCCAGGACACCTTGACC-3'	40	360
CD4	5'-GTA CTGAGGCAAGGTCCGCA-3' 5'-TGCACTATCTCCCAGAGCCA-3'	35	320
CD8	5'-ACCCTGAGAACTTCCGCCA-3' 5'-TGACCAATGAGAGGAGAAGG-3'	35	310
CD11b	5'-TCCTTTGCTGGGTGCTGTGG-3' 5'-GCAGGACAGCATCACACTTC-3'	35	440
CD45	5'-TTCTGCAACTGCTCCCTCGA-3' 5'-AATGGAGGCTCACATGAAGG-3'	35	240
G3PDH	5'-ATGGGGAAGGTGAAGGTCGG-3' 5'-ATCATATTGGAACATGTAAA-3'	35	150

Nucleotide sequence comparisons were performed using BLAST network program (National Center for Biotechnology Information, NIH, Bethesda, MD, USA).

Statistical Analysis

Light intensity (reverse transcriptase-polymerase chain reaction, RT-PCR) measurements were subjected to least squares (LS) ANOVA using the general linear model procedures of the Statistical Analysis System (version 6.0; SAS Institute, Cary, NC, USA). The model used in the LS-ANOVA included day (or treatment) and replicate as sources of variation. The light intensity from G3PDH PCR products was used as covariates for RT-PCR analyses. The least square mean (LSM) value and S.E. illustrated in the figures were derived from this analysis.

RESULTS

Expression of Various Cytokines in the Caprine Uterus

In agreement with our previous studies,^{24,25} IP-10 mRNA expression was higher in the endometrium of pregnant goats than that of cyclic goats (Fig. 1). The expression of IP-10 receptor CXCR3 mRNA also increased in endometrium of pregnant goats as compared with endometrium of cyclic goats. Levels of

IL-10, IFN- γ and TNF- α transcripts were also examined, and changes in IL-10 mRNA levels in endometrium of pregnant goats were found similar to those of IP-10 expression (Fig. 1). The presence of IFN- γ and TNF- α mRNA was detected in the endometrium of both cyclic and pregnant animals, but changes in mRNA levels of these cytokines were minimal.

Expression and Population of Immune Cells in the Caprine Uterus

To determine the distribution of immune cells in the pregnant uterus, the expression of CD4 (generally associated with Th lymphocytes), CD8 (usually associated with T cytotoxic lymphocytes), CD11b (NK cells and macrophages) and CD45 (all leukocytes) mRNA was determined using RT-PCR analysis. Levels of CD4 and CD11b mRNA were greater in endometrium of day 20 pregnant goats than endometrium from goats on other days examined; however, significant changes in CD8 and CD45 mRNA levels were not observed (Fig. 2).

Immunohistochemical analysis revealed that CD4, CD8 and CD11b positive cells were localized in cyclic and pregnant goat uteri (Fig. 3). Although a definitive conclusion could not be made, the number of CD4, CD8 and CD11b positive cells appeared to be higher in the subepithelial stroma and luminal epithelium of pregnant goats than those of cyclic goats. In addition, substantial numbers of positive cells were also

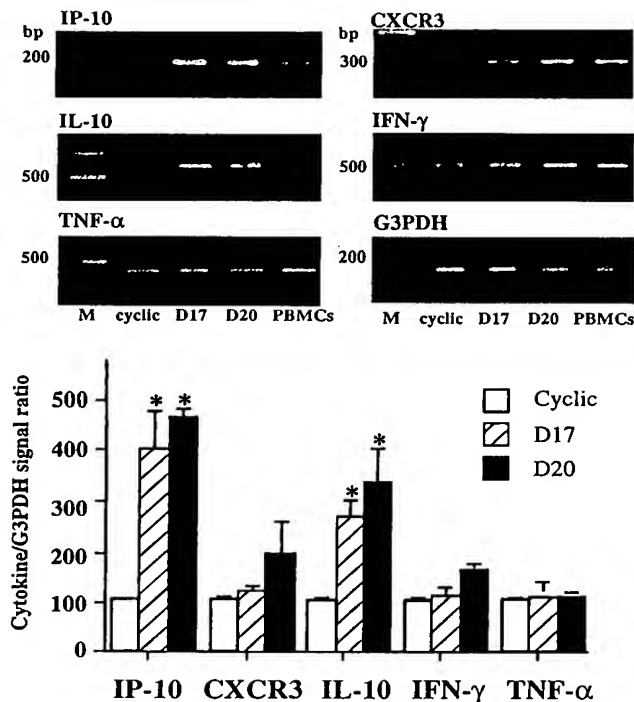


Fig. 1. Examination of chemokine and cytokine mRNA levels in the caprine endometrium. (Upper) RT-PCR analysis of changes in IP-10, CXCR3, IL-10, IFN- γ , TNF- α and G3PDH mRNA levels. Cyclic: day 15 cyclic goat endometrium ($n = 3$). D17 and D20: days 17 and 20 pregnant goat endometrium ($n = 3$ each). PBMCs: peripheral blood mononuclear cells from day 15 cyclic goats ($n = 3$). One set of representative data is shown. (Lower) Densitometric analysis of PCR products. Data represent those from three animals and each performed in triplicates. Bars represent LSM \pm S.E., and differences in these mRNA levels within a treatment group relative to that in cyclic animals are shown with an asterisk ($P < 0.05$).

localized on the surface of trophoctoderm in pregnant goats.²⁵ In the cyclic goat uterus, CD4, CD8 and CD11b positive cells were rarely localized in the subepithelial stroma (Fig. 3).

Effect of IP-10 on Immune Cell Migration

To ascertain the types of immune cells that migrated in response to the IP-10 treatment, isolated PBMCs were subjected to chemotaxis assay, followed by flow cytometric analysis against anti-CD4, anti-CD8 or anti-CD11b antibody. Chemotaxis assay showed that IP-10 increased the migration of PBMCs and the effect of IP-10 was neutralized by pre-treatment with the anti-IP-10 antibody (Fig. 4A). Flow cytometric analysis revealed that the population of CD4 and CD8 positive cells did not change in migrated cells stimulated with IP-10. But, the number of CD11b positive cell migration, when PBMCs were treated with IP-10, was higher than that of a group of cells without IP-10 stimulation. These data indicate that IP-10 could

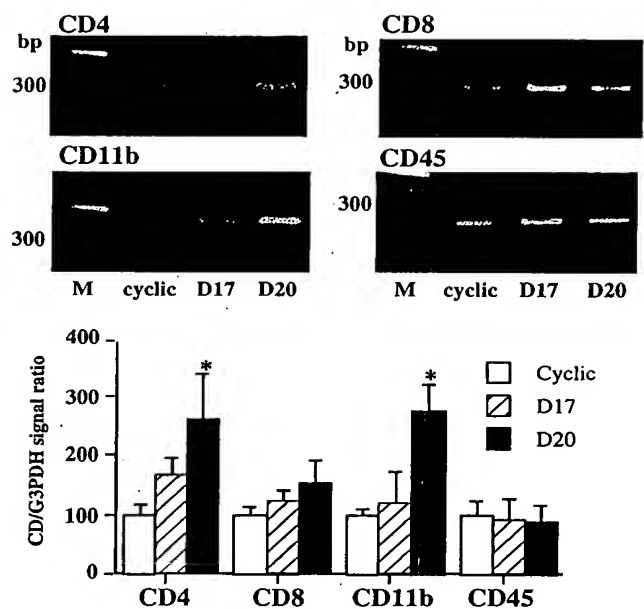


Fig. 2. Examination of CD4, CD8, CD11b and CD45 mRNA levels in the caprine endometrium. (Upper) RT-PCR analysis of expressions of CD4, CD8, CD11b and CD45 mRNA levels. Cyclic: endometrium from day 15 cyclic animals ($n = 3$). D17 and D20: endometrium from days 17 and 20 of pregnant animals ($n = 3$ each). One set of representative data is shown. (Lower) Densitometric analysis of PCR products. Data represent those from three animals and each performed in triplicates. Bars represent LSM \pm S.E., and differences in these mRNA levels relative to that in cyclic animals are shown with an asterisk ($P < 0.05$).

stimulate the migration of CD11b positive cells rather than CD4 and CD8 positive cells (Fig. 4B). The expression of CD4, CD8, CD11b and CXCR3 mRNA in migrated PBMCs with or without IP-10 stimulation were examined using RT-PCR analysis (Fig. 5A,B). It was confirmed that IP-10-stimulated migrated cells expressed more CD11b and CXCR3 mRNAs than those in non-stimulated migrated cells. However, the expression of CD4 and CD8 mRNAs did not change in migrated PBMCs with or without IP-10 stimulation (Fig. 5A,B). These observations were consistent with the results obtained from the flow cytometric analysis.

Effect of IP-10 on Cytokine Expressions

Cells that migrated with IP-10 stimulation had increased the expression of IL-10 and IFN- γ mRNAs, but the degree of increase in IL-10 mRNA was greater than that in IFN- γ mRNA (Fig. 5A,C). The expression of TNF- α did not change in migrated PBMCs with or without IP-10 stimulation. To determine whether IP-10 and/or IFN- γ could directly stimulate IL-10 expression, the expression of IL-10,

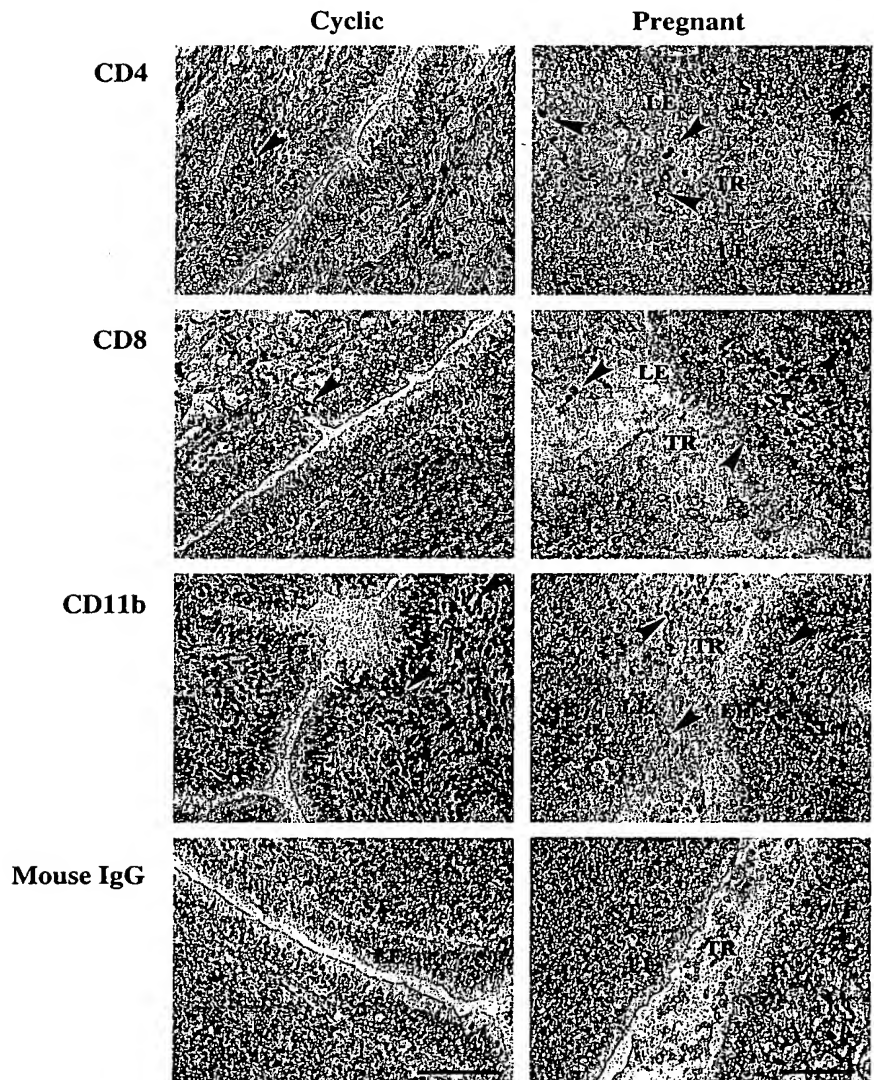


Fig. 3. Immunohistochemical detection of CD4, CD8 and CD11b in the cyclic (day 15) and pregnant (day 20) uteri. Antibodies used were anti-CD4, CD8 or CD11b mouse monoclonal antibody (all antibodies are cross-reactive to goats) and mouse IgG was used as a negative control. One set of representative data is shown. Arrowheads indicate positive signals. LE, luminal epithelium; ST, subepithelial stroma; TR, trophoblast. Bar = 100 μ m.

IFN- γ and TNF- α in PBMCs stimulated by IP-10, IFN- τ or recombinant GST protein (control) was investigated (Fig. 6). The low endogenous expression of IL-10 mRNA in PBMCs was observed using RT-PCR analysis (Figs 1 and 6A). IL-10 mRNA expression was stimulated with IP-10 or IFN- τ treatment, but the stimulation of IL-10 by IP-10 was greater than that by IFN- τ (Fig. 6B). The expression of IFN- γ mRNA was also stimulated by IP-10, but TNF- α mRNA was not affected by any treatments.

DISCUSSION

In addition to the prevention of luteolysis, proper interaction or adjustment between the conceptus and maternal immune system is required for pregnancy

establishment and maintenance. In ruminant ungulates, a conceptus factor IFN- τ is implicated in the process of maternal recognition of pregnancy by indirectly inhibiting CL regression.¹⁵⁻¹⁸ This IFN also regulates lymphocyte proliferation and cytokine production.¹⁹⁻²³ Recent studies revealed that endometrial IP-10 production appeared to be regulated by conceptus IFN- τ , and the migration of PBMCs was affected by IP-10 expression.^{24,25} Quite recently, the secretion of IP-10 and monokine induced by IFN- γ (MIG), both of which share the same receptor CXCR3, by cultured human endometrial stromal cells was found to be induced by progesterone, and endometrial IP-10 and MIG concentrations were correlated with the number of endometrial NK cells.³¹ Based on these observations, progesterone, conceptus IFN- τ and endometrial chemokines were suspected to play an important role in the

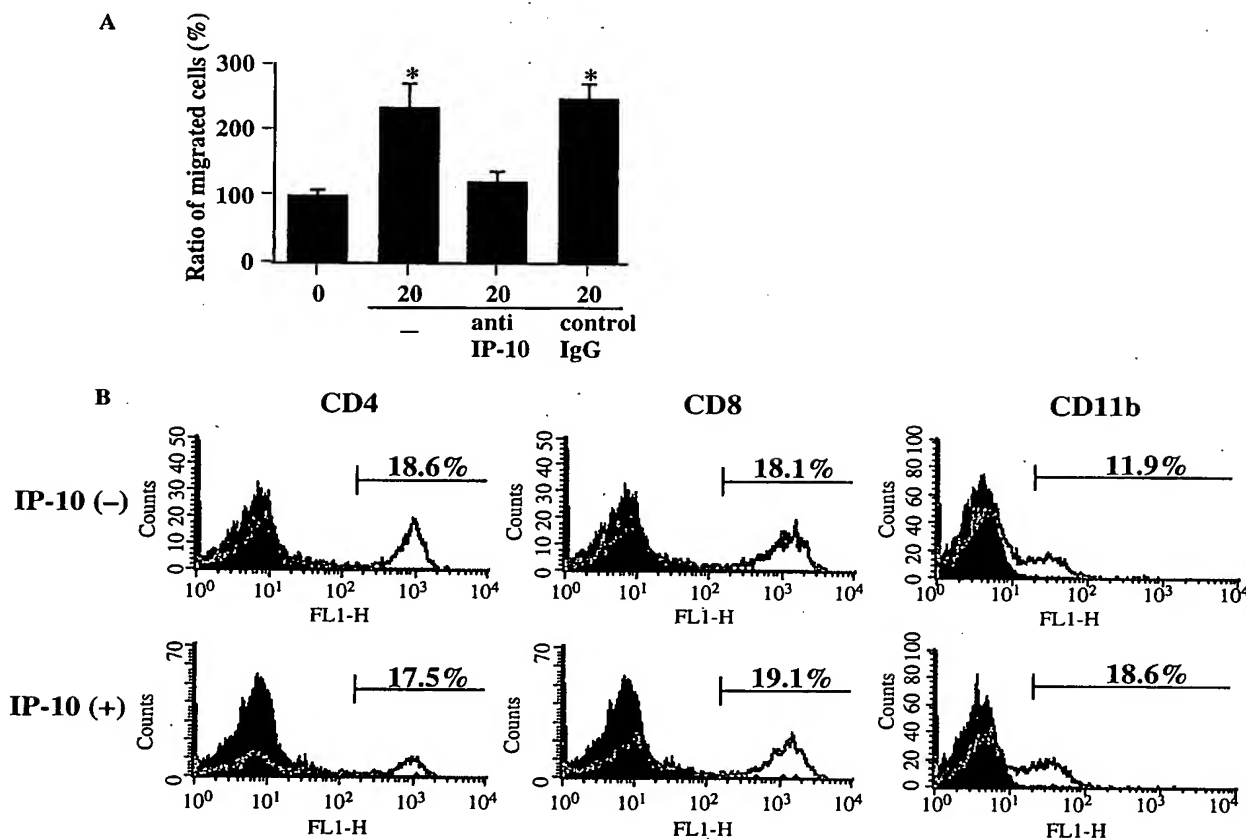


Fig. 4. Migration of PBMCs and changes in cell phenotypes with the IP-10 treatment. (A) Migration of PBMCs by IP-10. Effects of recombinant caprine IP-10 (20 ng/mL) on the migration of PBMCs pretreated with 30 μ g/mL of anti-IP-10 IgG or normal rabbit IgG were tested using a Boyden chamber. Bars represent LSM \pm S.E. An asterisk indicates a significant difference ($P < 0.05$). Three replicates were carried out for each treatment, and three independent experiments were performed for each treatment. (B) Expression of CD4, CD8 and CD11b cell markers on migrated PBMCs. Migrated cells in the presence or absence of IP-10 were harvested from three independent cultures. For flow cytometric (FACS) analysis, these cells were blocked against non-specific binding, and antibodies against CD4, CD8 and CD11b were then added to the cells. Cells reactive to cell marker antibody were identified using FITC-conjugated anti-mouse IgG. Numbers in figures indicate the percentage of positive cells. The experiment was performed three times independently.

immune cell distributions and patterns of cytokine productions at the endometrium, resulting in the adjustment and/or establishment of uterine environments suitable for the survival of ruminant conceptuses during early pregnancy.

In this study, the increase in IP-10 and IL-10 mRNA expression, but not IFN- γ or TNF- α was observed in the endometrium of pregnant goats. IL-10 is a cytokine with potent immunosuppressive effect,³² and is considered to play a crucial role in the alteration of immune-associated environments *in utero* during early pregnancy.³² It has been documented that regulatory cytokines including IL-10 are produced by Th2 lymphocytes and NK cells.^{10,11,33–36} In addition, increase in CD4 (generally associated with Th1 and Th2 lymphocytes) and CD11b (associated with macrophages and NK cells) mRNA expression was realized in the endometrium of pregnant goats. These results

indicated that endometrial IL-10 could be produced by Th2 lymphocytes and/or NK cells in the goat uterus. Furthermore, expressions of IP-10 and IL-10 in the pregnant endometrium appeared to be parallel, suggesting that endometrial IL-10 production could be increased with IP-10 stimulation. This notion was supported with the observations from chemotaxis assays in which PBMCs migrated with IP-10 stimulation had increased transcripts of IL-10, CXCR3 and CD11b, but not IFN- γ , TNF- α , CD4 or CD8. These results suggest that caprine IP-10 might affect migration or infiltration of NK cells rather than T lymphocytes or macrophages, which is consistent with the previous study in humans, in which IP-10 directed migration of NK cells by binding to and activating CXCR3 receptor.^{27,31,37}

It has long been suspected that uNK cells are required for the regulation of uterine environments in

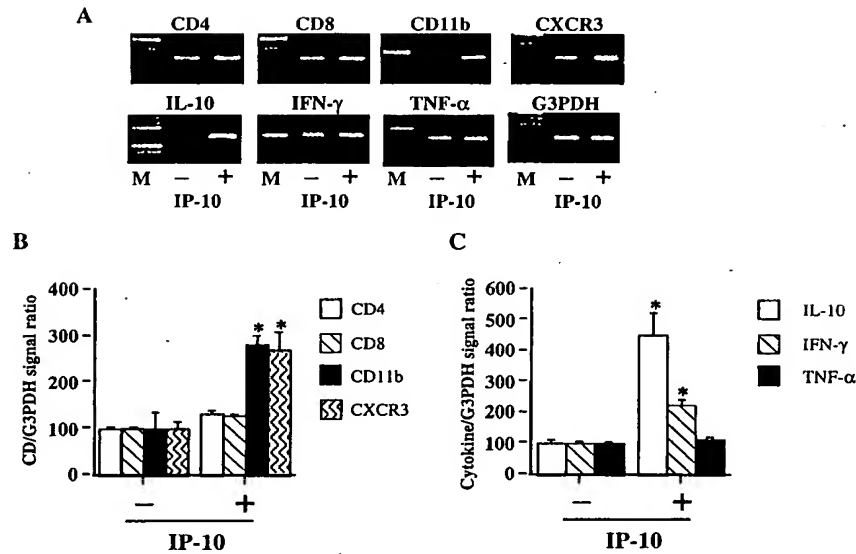


Fig. 5. Changes in cell surface marker and cytokine expressions on IP-10-stimulated migrated PBMCs or not migrated PBMCs. (A) Results from RT-PCR analysis for CD4, CD8, CD11b CXCR3, IL-10, IFN- γ , TNF- α and G3PDH mRNA levels on migrated PBMCs in the presence or absence of 20 ng/mL IP-10. One set of representative data is shown. (B) Densitometric analysis of RT-PCR products on CD4, CD8, CD11b cell surface markers, CXCR3 and G3PDH mRNA levels. Bars represent LSM \pm S.E., and differences in these mRNA levels relative to that of PBMCs without IP-10 treatment are shown with an asterisk ($P < 0.05$). (C) Densitometric analysis of transcript levels resulted from RT-PCR studies on IL-10, IFN- γ , TNF- α and G3PDH mRNA levels. Bars represent LSM \pm S.E., and differences in these mRNA levels relative to that of PBMCs without IP-10 treatment are shown with an asterisk ($P < 0.05$).

the human and murine pregnancy. When pregnancy occurs and advances, the number of uNK cells increases progressively. uNK cells are known to produce IFN- γ , which is involved in the local immunoregulation during early pregnancy.³⁸ In fact, the results from uNK cell-deficient mice reveal that IFN- γ concentrations at the implantation sites are considerably lower than those in wild-type mice, and deciduas of pregnant uterus exhibit necrosis, hypocellularity and abnormal vasculature.³⁸⁻⁴⁰ It is well documented that IFN- γ production is required for the establishment of murine pregnancy,³⁸ however, this proinflammatory cytokine is detrimental to the maintenance of pregnancy if the production persists beyond pregnancy establishment periods. Thus, the proper shift of the proinflammatory to regulatory cytokine productions by uNK cells may determine the success or failure of implantation processes. In ruminants, data related to uNK cell distribution and to IFN- γ production in the uterus are limited, but shifts of uterine environments detrimental to suitable for conceptus survival are required for pregnancy establishment.

In experiments with CBA \times DBA/2 mice (a spontaneous fetal resorption model), the intrauterine injection of recombinant ovine IFN- τ has reduced conceptus and embryo resorption.⁴¹ It is thought that increased production of uterine IL-10 by IFN- τ

administration is a reason for higher conceptus survival. In addition, intrauterine administration of anti-IFN- γ antibody to CBA \times DBA/2 mice reduces the embryo resorption rate.⁴¹ In the normal pregnancy model, CBA \times BALB/c mice, endometrial IL-10 production is much higher than that in the CBA \times DBA/2 mice, and when IL-10 is given to CBA \times DBA/2 mice, embryo and fetal mortality decreases significantly.⁴¹ The paradox in pregnancy immunology could be explained by one of the hypotheses that uNK cells in cyclic and pregnant endometrium produce both IFN- γ and IL-10, but when pregnancy occurs, uNK cells switch the cytokine production from IFN- γ to IL-10. In ruminants, it is possible that both conceptus factor IFN- τ and endometrial IP-10 have a role in the alteration of immune cell phenotypes toward uNK cells, which increase the production of IL-10 in the progesterone dominant-pregnant uterus.

In this study, IFN- τ as well as IP-10 stimulated the expression of IL-10 mRNA in PBMCs. The ability of IFN- τ to stimulate IL-10 production and to function as a immunosuppressor has been investigated previously.^{41,42} IFN- τ promoted the proinflammatory to the regulatory cytokine productions; however, biochemical and/or molecular mechanisms regulating such productions have not been clarified. Moreover, the exact mechanism by which IL-10 production is stimulated by IFN- τ *in vivo* has not been determined. Co-expression

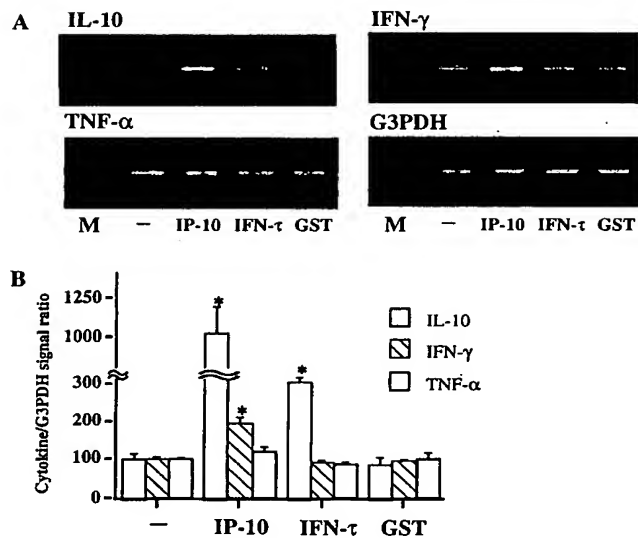


Fig. 6. Effect of IP-10 and IFN- τ on cytokine mRNA levels in PBMCs. (A) RT-PCR of IL-10, IFN- γ , TNF- α and G3PDH mRNA on PBMCs stimulated with 20 ng/mL IP-10, 10 ng/mL IFN- τ or control as 20 ng/mL recombinant glutathione S-transferase. One set of representative data is shown. (B) Densitometric analysis of RT-PCR products for several cytokines, IL-10, IFN- γ and TNF- α , mRNA and G3PDH mRNA. Bars represent LSM \pm S.E., and differences in these mRNA levels relative to that of PBMCs without any treatment are shown with an asterisk ($P < 0.05$).

and interaction of IP-10 and IL-10 has been reported from the studies of antitumor immunotherapy, which reveals that IP-10 is up-regulated in IL-10 expressing tumors and is important in the tumor-inhibitory activity of IL-10.^{43,44} In contrast, it was reported that IL-10 might act indirectly to suppress the expression of IP-10 from immune cells.^{45,46} Although the expression and interactions of IP-10 and IL-10 are demonstrated in many experimental models of anti-viral and antitumor effects and/or successful implantation, further investigations are required to elucidate effects and regulatory mechanisms of these factors during the period of pregnancy establishment in ruminant ungulates.

CONCLUSION

The results from the present investigation showed the expression of IP-10, IFN- γ , TNF- α , IL-10, CXCR3 mRNA and leukocytes cell surface markers, CD4, CD8, CD11b and CD45 mRNA in the caprine uterus. Data revealed that IP-10, found in the endometrium, could regulate the migration of PBMCs expressing IL-10 and CD11b, probably NK cells. These observations suggest that endometrial IP-10, regulated by IFN- τ and possibly progesterone, plays an important role in

the redistribution of uNK cells and/or the change of uNK cell phenotypes, which increase the production of IL-10. These changes in the local immune cell populations during the early pregnancy period play a crucial role for successful implantation in ruminants. Further investigations are required to better characterize and define a physiological role(s) for resident NK cells and NK cells recruited by endometrial IP-10 during early pregnant period in ruminants.

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